





590 600
CTTCAAGGAGGACGGCAACATCCTGGGGC
D05-Y75772-G0237014-4_G0237014-4-seq1r.ab1 (1>770) ← CTTCAAGGAGGACGGCAACATCCTGGGGC
G0237014-4.seq(1>869) → CTTCAAGGAGGACGGCAACATCCTGGGGC

610 620 630
ACAAGCTGGAGTACAACACTACAACAGCCAC
D05-Y75772-G0237014-4_G0237014-4-seq1r.ab1 (1>770) ← ACAAGCTGGAGTACAACACTACAACAGCCAC
G0237014-4.seq(1>869) → ACAAGCTGGAGTACAACACTACAACAGCCAC

640 650 660
AACGTCTATATCATGGCCGACAAGCAGAA
D05-Y75772-G0237014-4_G0237014-4-seq1r.ab1 (1>770) ← AACGTCTATATCATGGCCGACAAGCAGAA
G0237014-4.seq(1>869) → AACGTCTATATCATGGCCGACAAGCAGAA

670 680 690
GAACGGCATCAAGGTGAACTTCAAGATCC
D05-Y75772-G0237014-4_G0237014-4-seq1r.ab1 (1>770) ← GAACGGCATCAAGGTGAACTTCAAGATCC
G0237014-4.seq(1>869) → GAACGGCATCAAGGTGAACTTCAAGATCC

700 710 720
GCCACAACATCGAGGACGGCAGCGTGCAG
D05-Y75772-G0237014-4_G0237014-4-seq1r.ab1 (1>770) ← GCCACAACATCGAGGACGGCAGCGTGCAG
G0237014-4.seq(1>869) → GCCACAACATCGAGGACGGCAGCGTGCAG

730 740 750
CTCGCCGACCACTACCAGCAGAACACCCC
D05-Y75772-G0237014-4_G0237014-4-seq1r.ab1 (1>770) ← cTCGCCGACCACTACCAGCAGAACACCCC
G0237014-4.seq(1>869) → CTCGCCGACCACTACCAGCAGAACACCCC
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786) → CAGCAGAAcACCCC

760 770 780
CATCGGCGACGGCCCCGTGCTGCTGCCCC
D05-Y75772-G0237014-4_G0237014-4-seq1r.ab1 (1>770) ← CATCGGCGACGGCCCC
G0237014-4.seq(1>869) → CATCGGCGACGGCCCCGTGCTGCTGCCCC
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786) → CATCGGCGACGGCCCCGTGCTGCTGCCCC

790 800 810
ACAACCACTACCTGAGCTACCAGTCCGCC
G0237014-4.seq(1>869) → ACAACCACTACCTGAGCTACCAGTCCGCC
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786) → ACAACCACTACCTGAGCTACCAGTCCGCC

820 830 840
CTGAGCAAAGACCCCAACGAGAAGCGCGA
G0237014-4.seq(1>869) → CTGAGCAAAGACCCCAACGAGAAGCGCGA
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786) → CTGAGCAAAGACCCCAACGAGAAGCGCGA

850 860 870
TCACATGGTCCTGCTGGAGTTCGTGACCG
G0237014-4.seq(1>869) → TCACATGGTCCTGCTGGAGTTCGTGACCG
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786) → TCACATGGTCCTGCTGGAGTTCGTGACCG

Project: Untitled.SQD Contig 1

		880 890 CCGCCGGGATCACTCTCGGCATGGACGAG
G0237014-4.seq (1>869)	→	CCGCCGGGATCACTCTCGGCATGGACGAG
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	CCGCCGGGATCACTCTCGGCATGGACGAG
		900 910 920 CTGTACAAGTCCGGACTCAGATCTCGAGC
G0237014-4.seq (1>869)	→	CTGTACAAGTCCGGACTCAGATCTCGAGC
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	CTGTACAAGTCCGGACTCAGATCTCGAGC
		930 940 950 TCAAGCTTCGAATTCTGCAGTCGACGGTA
G0237014-4.seq (1>869)	→	TCAAGCTTCGAATTCTGCAGTCGACGGTA
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	TCAAGCTTCGAATTCTGCAGTCGACGGTA
		960 970 980 CCGCGGGCCCGGGATCATCAACAAGTTTG
G0237014-4.seq (1>869)	→	CCGCGGGCCCGGGATCATCAACAAGTTTG
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	CCGCGGGCCCGGGATCATCAACAAGTTTG
		990 1000 1010 TACAAAAAAGCTGAACGAGAAACGTAAAA
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	TACAAAAAAGCTGAACGAGAAACGTAAAA
		1020 1030 1040 TGATATAAATATCAATATATTAAATTAGA
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	TGATATAAATATCAATATATTAAATTAGA
		1050 1060 1070 TTTTGCATAAAAAACAGACTACATAATAC
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	TTTTGCATAAAAAACAGACTACATAATAC
		1080 1090 1100 TGTA AACACAACATATCCAGTCACTATG
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	TGTA AACACAACATATCCAGTCACTATG
		1110 1120 1130 GCGGCCGCATTAGGCACCCCAGGCTTTAC
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	GCGGCCGCATTAGGCACCCCAGGCTTTAC
		1140 1150 1160 ACTTTATGCTTCCGGCTCGTATAATGTGT
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	ACTTTATGCTTCCGGCTCGTATAATGTGT
		1170 1180 GGATTTT GAGTTAGGATCCGTCGAGATTT
G11-Y75772-G0237014-4_G0237014-4-seq2.ab1 (1>786)	→	GGATTTT GAGTTAGGATCCGTCGAGATTT

